## PPTENE ITAMIN

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+ [. + PINISHE FERTIFIC FOR CARCINOMMERYODIC ANTIGEN (CEA.
-156 - Sequence Listing DVX-016.6 US
Pitana MD 00/841,841
21419 2000-54-63
<!!!> 137
<216 % 1
<211 - 16
<212 - PRT
<213 Artificial Sequence
 23. Description of Artificial Sequence: CEA binding
      polypeptide
<220:
<2211 VARIANT
₹222. (1)
4223 Kaa is Asn, Asp or is absent
12305
<221. VARIANT
12/22: [2]
12:13 Xaa is Trp
12.20.
:221 VARIANT
<222:- (3)
<223: Xaa is Asp, Phe or Val
<221: VARIANT
42221 (5)
<223 Xaa is Ash, Glu or Met
-2211 VARIANT
42.52 164
<223 Maa is Ass, Lew, Met or Phe
    VAFIANI
this Kaa is Asy, Gly, The, Lys, Pho or Thr
<2210 VARIANT
12222 (5)
<fi><fi>> Xaa is Ala, Gir, Gly, Lys or Thr
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Booker Kaarook erd, Albo, Aug, Albord by
Total Conference
RAABR Maa is Bin, Leutor Gly
42261
<2.21> VARIANT
*8722 1117
<233> Xaa is Ala, Trp or Tyr
<2205
<221: VARIANT
42221 (12)
<223 - Kaa is Ala, Gly, His, Phe, Thr or Val</p>
KARTANT
<2.22: (14)
<2230 Xaa is Asn, Gln, Phe, Ser or Val
<2201-
<221: VARIANT
-222- (15)
<223> Xaa is Arg, Leu, Pro or Ser
<220.
GAZI, WARTANT
<722- (16)
/223: Maa is Leu, Ser, Trp or Tyr
<4001-1
Kaa Xaa Kaa Cys Kaa Kaa Kaa Kaa Kaa Kaa Kaa Kaa Cys Kaa Kaa Kaa
                                    10
1210: 2
<2111 16
Ollah PRT
Wille Artificial Sequence
1.25 Description of Artificial Sequence: family of
     preferred CEA binding moieties
(%%3> Kaa is Asn or Asp
<221> VARIANT
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· ... · · ·
 over Maarin Hime, Met, Detre fr Akh
- Lik variasi
+ 1. W. Maa is Asr, dly, lle, lys, the or Thr
· // LZ VARIANT
· 1.2.2 9.
2008 Maa is Arg, Asn, Asp, Glu, Gly or Trp
12.291
<221 VARIANI
<222: (12)
<223: Xaa is Ala, Gly, His, Phe, Thr, Tyr or Val
r.220%
 GERTE VARIANT
 1222 (11)
 ...3 - Måa is Arg, Leu, Pro or Ser
4.2201
<221: VARIANT</pre>
42221 (16)
<223: Xaa is Leu, Ser, Trp or Tyr
100. 2
Maa Trp Val Cys Glu Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Xaa Xaa
                                        10
*216. 3
4,211, 10
<212: PRT
<213: Artificial Sequence
<220.4
<223: Description of Artificial Sequence: CEA binding
      loup
· 220,
• 231,
• 112, ×
      VARIANT
*213 Kaa is Asm, Glu or Met
 ) i, VARIANT
ina (3)
 of Eas is Asi, Log, Mot or Phe
KSSI - MARIANT
~222 . . . 41
~223> Waa is Asp, Gly, Ile, Lys, Phe or Thr
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on to Haa for Ana, Ann, Hy, Lys Or Thr
* X.1X.* (6)
- 753 - Maa la Arq, Ash, Asp, Giu or Gly
<121 VARIANT
< 222.
<123: Maa is Glm, Gly or Leu
<221. VARIANT
<2221 (8)
<223: Xaa is Ala, Trp or Tyr
WERE VARIANT
-822: :91
<2230 Xaa is Ala, Gly, His, Phe, Thr or Val
<4001 3
Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys
<210:-4</pre>
s.2111-- 16
~212.- PR4
<213. Artificial Sequence</pre>
<223: Description of Artificial Sequence: CEA binding
    polypeptide
<400: 4
Asr. Tro Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asn Ser Tyr
                            10
 1
.115> 5
47112 16
PRIJA PRI
20132 Artificial Sequence
- (3) Description of Artificial Sequence: CEA binding
     -palymeptide
Asy Try Val Tys Glu Ash Lys Lys Asp Gln Trp Thr Cys Ash Leu Leu
                                    10
```

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The Artificial Corporations
+...lar beseription of Artificial Sequence: CFA binding
    polypeptiae
1415> 6
Asn Trp Asp Cys Met Phe Gly Ala Glu Gly Trp Ala Cys Ser Pro Trp
<210> 7
<211> 16
ィ2キラス PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
    polypeptide
<400> 7
 Asp Trp Val Cys Glu Lys Thr Thr Gly Gly Tyr Val Cys Gln Pro Leu
                              10
 1 5
 <210> 8
 <211> 16
 <2120 PRT
 <213> Artificial Sequence
 222C>
 <223> Description of Artificial Sequence: CEA binding
     polypeptide
 Asn Trp Phe Cys Glu Met Ile Gly Arg Gln Trp Gly Cys Val Pro Ser
                                  10
 +111.x 3
×211> 16
  2012× PRW
 · 21 to Artificia, Sequence
 polyportido
  4175 3
  Asy Trp Val Tys Asn Phe Asp Gln Gly Leu Ala His Cys Phe Pro Ser
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e. Ire Artificial Sequences
domain for design of migroprotein display library
PARIA VARIANT
-122> (1)...(12
<2235 amino acid positions 4 and 9 are invariant Cys;
     all other positions Xaa are varied but not Cys, to
     provide a library of 2x10(8) different peptides
     based on the template sequence
<410-10
Yaa laa Kaa Cys Maa Xaa Xaa Maa Cys Xaa Xaa Xaa
<210: 11
<211: 11
<212: PRT
213: Artificial Sequence
4223: Description of Artificial Sequence: parental
     domain for design of microprotein display library
<221: VARIANT
<222 (1)..(11)
#223. amino acid positions 3 and 9 are invariant Cys;
     all other positions Xaa are varied but not Cys, to
     provide a library of 1x10(3) different peptides
     based on the template sequence
<400.- 11
Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
                                   10
F. 10: 12.
-XIX Artificia, Coquence
Alair Pestription of Artificial Sequence: parental
     demain for design of microprotein display library
KIZI - DARTANT
POSTA aminu and positions 3 and 10 are invariant Sys;
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     gar vides a lightary of TMI? Politicement perhips.
     pared in the template bedrets
Maa Maa Cyn Maa Maa Maa Maa Maa Cys Maa Maa
10
42.10% 13
77:13 16
Aliza PET
7213> Artificial Sequence
<2200
<223: Description of Artificial Sequence: parental
      domain for design of microprotein display library
x $ 21 - WAR IANT
<222 - 10) ... (16)
%223! amino acid positions 4 and 13 are invariant Cys;
      all other positions Xaa are varied but not Cys, to
      provide a library of 2.5x10(8) different peptides
      based on the template sequence
<400 13
Xaa kaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa
                                    10
 1 5
3210 - 14
< 211 - 16
<212 - PRT
<213 · Artificial Sequence
<220 ·
<223> Description of Artificial Sequence: variable
     sublibrary sequence used in designing focused
      secondary library
<220>
 <221> WARTANT
 2007 Mad is any amino acid except Cys
 -121> VARIANT
 ∠2.22 > (5)..(6)
 · (2) A Kaa is any amino acid except Cys
 -17.2 14
 Kaa Kaa Kaa Cys Kaa Kaa Lys Lys Asp Gln Trp Thr Cys Asn Leu Leu
                                     10
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all the selection of the present the
. . . .
. . . . . . . . . . . . . . . Artificial Sequence: variable
     mur. Extery to pier to used in designing focused
     segmnasy library
77210 TARIANT
·222> (B)...9

XAABS Xaa is any amino acid except Cys
Asp Irp Val Cys Xaa Xaa Xaa Xaa Gln Trp Thr Cys Asn Leu Leu
-7.162-16
-2115-16
<212> PRT
<213: Artificial Sequence
<220:
<223: Description of Artificial Sequence: variable
       sublibrary sequence used in designing focused
       secondary library
 <220 •
 <221 · VARIANT
 x222 · (8) .. (42)
223 · Xaa is any amino acid except Cys
 <400 - 16
 Asp Tro Val Cys Glu Asn Lys Xaa Xaa Xaa Xaa Xaa Cys Asn Leu Leu
                                       10
 <210 - 17
 <211 > 16
 <212> PRT
 <2135 Artificial Sequence
 ಳು. ಎ Description of Artificial Sequence: variable
        surflictory sequence used in designing focused
        secondary library
 RELIGION MARIANT
 21.222 (11) ... 12
 <223> Xaa is any amino acid except Cys
  POZIP TARTATT
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Ang lig Wal trye top Asso light lys Asp. Gls. Man Kan Tys Kan Kan Kan
                                           10
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- / 122 PET
82132 Artifitial Sequence
- 21.95
<223> Description of Artificial Sequence: variable
       sublibrary sequence used in designing focused
       secondary library
 ZZELZ ZARIANT
 e0220 6)..(7)
<2230 Xaa is any amino acid except Cys</pre>
 <2201
 <2211 VAF.IANT
 <2221 (9)
 <223 - Kaa is any amino acid except Cys
 <220>
 <221> VARIANT
  <222> (12)
  <223> Xaa is any amino acid except Cys
 <2205
  <221> VARIANT
  < 222 > (15)
  <2230 Xaa is any amino acid except Cys
  <400> 18
  Asp Irp Val Cys Glu Xaa Xaa Lys Xaa Gln Trp Xaa Cys Asn Xaa Leu
  <210> 19
   @11> 16
   3012> PFT
   p. so Artiflesal Sequence
  Fig. Description of Artificial Sequence: variable
         sublibrary sequence used in designing focused
         secondary Tibrary
   effic VARIANT
   Fall & 5 ...
   47775 Maa is any arino acid except Sys
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the site Mara 12 applications artificial six wept of the
  The trace is any aming used except bys
 1 4 1 VX 1 1
 Ast. Trp Val Cys Kaa Xaa Xaa Lys Xaa Gin Trp Xaa Cys Asn Ser Tyr
 <210> 20
 <211> 16
 <212> PRT
 <213> Artificial Sequence
 <923> Description of Artificial Sequence: variable
       sublibrary sequence used in designing focused
       secondary library
 <220>
 <221 - VARIANT
 <2221 (1)
<2230 Xaa is any amino acid except Cys
<2200
<221: VARIANT
<2222 (3)
<223) Xaa is any amino acid except Cys
<2200
<221> VARIANT
<222≥ (14)..(16)
<223> Xaa is any amino acid except Cys
<400> 20
Kaa Trp Kaa Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Kaa Kaa Kaa
                                      10
-210/ 21
· 711 * 16
-112 - FRI
n ing Artificia. Sequence
The st Description of Artificial Sequence: isolate of
      TN10/9 library found not to bind CEA
74009 21
Ash Trp Arg Dys Lys Leu Phe Pro Arg Tyr Pro Tyr Cys Ser Ser Trp
                                      10
```

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outo Artifica de penee
 to be bestription of Artificial Sequence: isolate of
      THIS A . Herary tound not to bind CEA
 Ar: Tyr Cys Stu the Phe Pro Trp Ser Leu His Cys Sly Arg Pro
                                     10
 <210> 23
 <211> 16
 <212> PRT
<213> Artificial Sequence
1.241
<2232 Description of Artificial Sequence: conserved
      amino acid positions in first family of CEA
      binding peptides
<220 ⋅
<221 · VARIANT
<222 (6)
<223 X is Asn, Leu, Met or Phe
<220
<221 VARIANT
3222. (7)
$223: X is Asp, Gly, Ile, Lys, Phe or Thr
42200-
<221: VARIANT
<222: (9)
<2230 X is Arg, Asn, Asp, Glu or Gly
<22200
<221> VARIANT
<222> (12)
<223> X is Ala, Gly, His, Phe, Thr or Val
HARIANT
ಳೂತಿ X is Arg, Lou, Pro or Ser
Asp lip Wal Dys Blu Maa Maa Lys Maa Cln Trp Maa Cys Ash Maa Leu
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 to the lest digits, hour Antificial Geopher were myminetre day.
       finding pertiae with "-terminal immobilization
        Beduction,
 *4". + 2.4"
 the Ash irr Mai Mys Ash Leu Phe Lys Ash Gin Trp Phe Cys Ash Sor
  Tyr A.a Pro Sly Gly Glu Gly Gly Gly Ser Lys
20 25
 <210> 25
 <211> 27
 42123 PRT
 *2132 Artificial Sequence
 <223> Description of Artificial Sequence: synthetic CEA
       pinding peptide with C-terminal immobilization
       sequence
 <400> 25
 Ser Asp Trp Val Cys Glu Asn Lys Lys Asp Gln Trp Thr Cys Asn Leu
 Leu Ala Pro Gly Gly Glu Gly Gly Gly Ser Lys
<210> 26
<211> 27
<212> PRT
<213> Artificial Sequence
<2205
<223> Description of Artificial Sequence: synthetic CEA
      binding peptide with C-terminal immobilization
      sequence
<406> 2€
Sor Asn Trp Asp Cys Met Phe Gly Ala Glu Gly Tip Ala Cys Sor Pro
Tip Ala Fro Siy Siy Giu Giy Giy Giy Ser Lys
· 2.11> 27
e2115 27
12129 FRT
Aliaz Artificial Sequence
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The to the trapts of Artific that dequences synthetic float
                       and the property of the second section of the second of th
        be Aug Day Tull by All be the The Gly Gly Tyr Val Cys Gin Fro 10 ^{\circ}
       Low Ala Pro Mily Gly Glu Gly Gly Gly Ser Lys 25
      4.316> 28
      <211> 10
      <212> PRT

Artificial Sequence
     C221.3
    <223> Description of Artificial Sequence: C-terminal
                         sequence for immobilizing poptides
     <4905 28
    Ala Pro Gly Gly Glu Gly Gly Gly Ser Lys
   <2101-29
   <211. 16
    <212: PRT
   <213: Artificial Sequence
   <2233 Description of Artificial Sequence: template
                      sequence for sublibrary used in construction of
                      focused secondary display library
  <220×
  <221> VARIANT
  <222> (1)..(3)
 <223> X is any amino acid except Cys
 ≺ŻŹI> VARIANT
 52428 (B ...(6)
 <223> X is any amino acid except Cys
-4738 29
 Maa Kaa Cys Kaa Kaa Lys Lys Asp Gln Trp Thr Cys Ash Leu Leu
                                                                                                                             10
R0112 30
<211> 16
Küllə PPT
12132 Artiliqua. Sequençe
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        in the world of the large large and the element become a
        d fined by hadry duglay a grary
 ozie vakláni
 +25.18~{
m K} is any amino acid except Cys
  Asp Trp Val Cys Edd Kad Kad Kad Kad Gln Trp Thr Cys Asn Leu Leu
 <210> 31
 <211> 16
 <212: FRT
 <213> Artificial Sequence
 12255
 <223> Description of Artificial Sequence: template
       sequence for sublibrary used in construction of
       focused secondary display library
 <220 ⋅
 <221 - VARIANT
 <222 - (8)..(12)
 <223: X is any amino acid except Cys
 K44002 31
Asp Trp Val Cys Glu Ash Lys Xaa Xaa Xaa Xaa Xaa Cys Ash Leu Leu
                                       10
<210> 32
<211> 16
<212> PRT
<213> Artificial Sequence
<22Gb
<223> Description of Artificial Sequence: template
      sequence for sublibrary used in construction of
      focused secondary display library
<220>
<221> VARIANT
· (172 × '11) . . (12)
Parties X is any action acid except Cys
· 601. - 27.18 7.1.
12:28 .14, ... 16
<223> X is any amino acid except Cys
14112 32
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og lid van ly lid Am lye lye Asp Nin Kam Maa Cy. Kam Kam Kam
  Tailan Artists that the paging
  Filis Lescription o: Artificial Sequence: template
        sequence for sublibrary used in construction of
        focused secondary display library
 82657
  #2212 VARIANT
  <222> (6)..(7)
 <222^{\circ}> X is any amino acid except Cys
 <221 >
 <221 - VARIANT
 €222 (9)
 \stackrel{\textstyle <}{\scriptstyle \sim} 222 \cdot \stackrel{\textstyle \times}{\scriptstyle \times} is any amino acid except Cys
 3220 .
 <221 - VARIANT
 <222 (12)
 <\!223\cdot\,X is any amino acid except Cys
 <221 · VARIANT
 <222 - (15)
 <223 - X is any amino acid except Cys
 <450 - 33
 Asp Trp Val Cys Glu Maa Maa Lys Maa Gln Trp Maa Cys Asn Maa Leu
<2107 34
<211: 16
<212: PRT
<213 Artificial Sequence
52.600
<
       sequence for sublibrary used in construction of
       focused secondary display library
+721× VARIĀNI
tivity H is any amino abid except Cys
. 22.17
<271> VARIANT
11,727 3.
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the trib in any amin actionex rept mys
 Ash Tro Va. Tys Mil Mod Maa Lys Maa Glo Tro Mad Cys Ash Der Tyr
P2112 16
PAILS PRI
8213> Artificial Sequence
1220>
<223> Description of Artificial Sequence: template
      sequence for sublibrary used in construction of
       focused secondary display library
#221 · WARIANT
4822 • (1)
<223 · X is any amino acid except Cys</pre>
<221: VARIANT
42220 (3)
\!<\!223\!\times X is any amino acid except Cys
72210 VARIANT
4%22) 114,...(16)
<223% X is any amino acid except Cys
<400> 35
Maa Trp Xaa Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Xaa Xaa Xaa
<210> 36
×211> 16
មាន 1.2 ២ ខេត្ត
<2132 Artificial Sequence
전23> Description of Artificial Sequence: family of CEA
      binding polypeptides
Burning TARIANT
*121> Maa is Asp, Ash, Ala or Ile
¥ 1.1 ° 3
```

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talite Kaaraa Caa, Law, Wet, Tyr, Ene, Ers er Aug
Bunger Kala ili Aim, Guush Asp
<121> VARIĀNI
€2.2.2> 16;
<223> Xaa is Leu, Phe, Typ, Trp, Val Met, Ile or Asn
-2200
Pariant JARIANT
<2225 (T)
*223> Kaa is Phe, Leu, Asp, Glu, Ala, Ile, Lys, Asn,
      Ser, Val, Trp or Tyr
*220%
<221> VARIANT
<222> 18)
<2235 Kaa is Lys, Phe, Asp, Gly, Leu, Asn or Trp
<220.
KOSTO VARIANT
<2220 (9)
<223> Xaa is Asn, Pro, Phe, Gly, Asp, Ala, Ser, Glu, Gln
      or Trp
<2200
₩221: VARIANT
$2225 (10)
ชน์นิธิภ Xaa is Glr or Lys
*2205
3221> WARIANT
₹222> (12)
<223> Xaa is Phe, Thr, Met, Ser, Ala, Asn, Val, His,
      Ilé, Pro, Trp or Tyr
<220>
RAPIN VARIANT
₹222> (14)
(223) Kaa is Asn, Asp, Glu, Pro, Gln or Ser
+221= WARIANT
12222 (15)
223 Xaa is Val, Leu, He, Pro, Ala, Gln, Ser, Met,
      Hu, Thr, Lys or Trp
42212 VARIANT
11112 16
PIFFE Maa is Leu, Mat, Val, Tyr, Ala, Fle, Trp, His,
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ar , éan, e., iné, ist a Afri
Kalandep Karanda, Mara Kara Kara Kara Kara Karandep Karandya Kara Karandara
enlige Artifical is graft se
+2230 Description of Artificial Sequence: CEA binding
    polypeptiae
4 3
Asy Trp Mot Tys Ash Leu Fhe Lys Ash Glh Trp Phe Cys Asp Leu Met
·..1... 38
·..11... 16
·!!!!> PPT
+113> Artificial Sequence
1 1.112
+223% Description of Artificial Sequence: CEA binding
      polypertide
R400> 38
Ask Trp Val Cys Ash Leu Phe Lys Ash Gln Trp Phe Cys Asp Leu Met
                         10
... 2 33
... 1: 2 1:
... 1: 2 PRT
Artificial Sequence
*** Description of Artitibial Sequence: CEA binding
      pulypoptide
A 11 (18) (54)
Asy Tip II: Tys Ash low the Lys Ash Glr Trp Phe Cys Asp Glr Mct
- 1 . 4
-.110 1.
- 11 . FMT
P. In Artistia, Popular
-.i.x
-firx Description of Arthiridal Sequence: CEA binding
      31 . 73 + 1 · . 4.
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and legal to the engineering to the System continuing the System and the
rout Artigolia, Tequality
-1.3 - Les maption of Artificial Dequence: CEA binding
     polypertide
14 m > 41
Asy Trp lie Cyc Ash Leu Phe Lys Ash Gln Trp Phe Cys Gln Val Lys
· ...Ze PKI
<213> Artificial Sequence
FANGE Description of Artificial Sequence: CEA binding
     - prlyrentlde
1 4 52 15
Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Met
*!:13 - Artificial Requestor
<223> Description of Artificial Sequence: CEA binding
     polypeptide
< 11 - 43</p>
Asy Try Mat Cys Ash Lau Phe Lys Ash Gln Try Phe Cys Asp Gln Fle
+.... + 44
+.... + 16
+.... + 18T
e ter Artificia, sequence
Music Legislich is Artificial Sequence: CEA binding
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 *
  -112 - IPT
 ellik Artifisial Engueroe
 Calibe Description of Artificial Sequence: CEA binding
       polypopinge
 <4(0> 45
 Asp rp lie Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Ile Arg
                       10
 <210: 46
 4211× 16
 Kalar PRT
 📆 13: Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: CEA binding
      polypeptide
 <400> 46
 Asp Trp Met Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Val
                5
                                   10
8710> 47
<211> 16
73132 PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 47
Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Ala Ilo
                            10
, bir-- 15
· 2115 16
1212> IKT
<213> Artifitial Sequence
+223 * Description of Artificial Sequence: CEA binding
     polypertide
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  talit Artiflera Gespense
  HART'S Les miption of Artificial Sequence: CEA binding
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  <4665 49
  Asp Trp Val Cys Glu Phe Leu Lys Met Gln Trp Ala Cys Asn Val Deu
 ## 11× 16
 -212× FRT
 หมิโร๊> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: CEA binding
       polypeptiae
 <400> 50
 Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asn Val Met.
                 5
 <210> 51
 <211> 16
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 <213> Artificial Sequence
5223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 51
Ala Trp Pro Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Pro Pro Gln
1. 1. 141
ा । वर्ष Artificial Sequence
<223> Description of Artificial Sequence: CEA binding
      polypeptide
the Tie Tal Tys Ash Leu Fre Lys Ash Gln Trp Pho Cys Asp Val Leu
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+. 11 + 10
+. 1. + EFT
  olite Artifologo Despende
 +12% + Les martler et Artificial Sequence: CEA binding
      pslypopilde
 化有货机区 1.3
 Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Lys Trp
                         10
 8210> 54
 <111× 16
<112× 287
/2:3> Artitinial Sequence
<2205
<223 \rightarrow Description of Artificial Sequence: CEA binding
      polypeptide
<400> 54
Asp Trp Val Cys Glu Trp Leu Lys Met Gln Trp Ala Cys Asn Met Leu
                                      10
<210 - 55
<211: 16
<212: PRT
<213: Artificial Sequence
<2200
<223> Description of Artificial Sequence: CEA binding
      polypoptide
<4035 55
Asp Trp Val Cys Asp Phe Phe Phe Asn Gln Trp Thr Cys Asn Leu Leu
<210× 56
<211> 16
5215 × 11:5
+%13× Ārtīfi da. Sequeñoc
ATTIC Les ription of Artificial Sequence: CEA binding
     polypeptide
<41' > 56
Asp Trp Val Cys Glu Met Phe Lys Ala Gin Trp Phe Cys Ash Ala Lep
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    p. Beschipt, here of Artificial Sequence: CEA binding

     aslypertiae
2 4 1 2 2 2 E
Asp Irp Ise Cys Ash Leu Phe Lys Ash Gln Trp Phe Cys Asp Ala Trp
                                     10
<210> 58
42112-16
-1125 PRT
e.ls Artifictal Sequence
1.25.
<223> Description of Artificial Sequence: CEA binding
     polypeptide
<400> 58
Asp 1rp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Trp
                                     10
<210: 89
87113 16
THILL PRI
<213> Artificia: Sequence
< 220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400: 59
Asp Irp Val Cys Glu Tyr Phe Lys Asn Gln Trp Phe Cys Asn Val Leu
                                     10
+211+60
<7111 16
KATAN PRI
+213 - Artificial Sequence
   - - Pastription of Artificial Sequence: CEA binding
     - polypeștidă
<4002 60
Asp Trp Val Cys Glu Ile Asp Lys Gly Gln Trp Thr Cys Asn Pro Leu
```

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taana les miptist of Artificial Sequence: CEA binding
    godypeptico
Asp Trp Ta. Dys Ash Leu Phe Lys Ash Gin Trp Phe Cys Ash Pro Phe
      B 10
<210⊳ 62
<211:- 16
<212: PRT
<213: Artificial Sequence
and a Description of Artificial Sequence: CEA binding
     polypeptide
Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Val Gln
<210: 63
42110 16
RAIDS PRT
<21:0 Artitimial Sequence
<223> Description of Artificial Sequence: CEA binding
     polypeptide
<400> 63
Asp Trp Val Cys Asn Leu Phe Phe Gly Gln Trp Thr Cys Asn Leu Leu
                   10
RA139-64
⊈211> 1€
-1155 PRT
-%id> Artif:dial Sequence
Al. to Description of Artificial Sequence: CEA binding
    pšlypapilin
Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Glu Ala His
```

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tamit lydstriptish 6: Artifilias Dequence: ChA binding
     polypeptien
 · 1 2 > 6.5
 Amp Trp Val Cys Glu Leu Val Lys Ala Gln Trp Tyr Cys Asn Ile Leu
                                     10
42152 66
 <211> 16
 <212> PRT
<2135 Artificial Sequence
Alian Description of Artificial Sequence: CEA binding
      pblypeptide.
4400× 66
Asn Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Thr Val
                             10
<210> 67
<211> 16
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: CEA binding
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<400> 67
Asp Trp Val Cys Glu Phe Tyr Lys Ser Gln Trp Asn Cys Asn Ile Leu
                                    10
<210> 68
<211> 16
4212> PRT
4913> Artificial Sequence
+1532 Description of Artificial Sequence: CEA binding
     polypertide
- i . Fi
Ask Trp Wal Cys 3.2 Trp Fhe Lys Pro Gln Trp Phe Cys Asn Pro Leu
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the traces digital bed estata del Despenses KNA binding
      Franklin in Pre-
· 第一、原传
her irr lyr lys Ash Leu Phe Lys Ash Gin Trp Phe Cys Ash Val Leu
92105 76
4211> 16
<212> PRT
<213> Artificial Sequence
<220>
(253) Description of Artificial Sequence: CEA binding
     relypoptiae
<4005 70
Asp Trp Val Cys Glu Tyr Asn Asp Glu Gln Trp Thr Cys Asn Leu Leu
                                     10
<2165 71
<211: 16
<212. PRT
<213: Artificial Sequence
<223. Description of Artificial Sequence: CEA binding
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<450: 71
Asp Trp Ile Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asn Glu Ala
<2185 72
<211> 16
<212> PRT
<213> Artificial Sequence
4 4.2 3.3
12.33 Description of Artificial Sequence: CEA binding
     polypeptide
Ag Iff Mai Tys Ash Trp Glu Leu Phe Gln Trp Thr Cys Ash Lou Leu
42132 73
82115 16
```

1 1 1 1

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e versie engitu not Artifolio Majaenser Waddinaing
wig ling Tall Tyr Arm Lott Ene Liys Arm Sain Tro, Ehê Cys Arp Sain Val
+..11;- 74
+2115+ 16
12.12.8 FRI
RMI>> Artifly al Sequence
127
<223> Description of Artificial Sequence: CEA binding
     polypoptida
. · · ·
Asy Inc Ta. Typ Ash Let Ene Lys Ash Glin Tro Phe Cys Asp Val Pro
<2.105 75
·211> 16
+2.12> PRT
Pulta Artificial Sequence
1.10 Description of Artificial Sequence: TEA pinding
     polycopt an
+46 0 Tb
Asp Irp Val Cys @lu Fhe Phe Lys Gln Gln Trp Phe Cys Asn Val Leu
                                 10
+ 111 1r.
+ 113 Hell
H. IHA Artiffe Fall Segret re
1.24 Description of Artificial Sequence: CEA binding
     - polypantide
Asy the transfer the fine the type Asy Bin Try Cor Cys Ash Wallier
/ 1 / 1 *
/...11* 16
×utub ≘ĒI
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4.5 / 14
4.55% 56
4.5. / 1.4
7. 1 w Artuit mai Sequence

A// -> Description of Artificial Sequence: CEA binding
     polypeptide
Aug Ing Tal Tys Glu Phe Met Lys His Gln Trp Phe Cys Asn Pro Leu
                                      10
3,17× 79
//.11> 16
k/12> PRT
दरी विशेष Artificial Sequence
්ධ්යා Description of Artificial Sequence: CEA binding
     prlyphrtian
Asp irp lie Cys Ash Leu Phe Lys Ash Gln Trp Phe Cys Gln Ala Val
          1.0
*2112 67
*211= 16
-711 PRT
* 1. * Artiblola : daquence
-1.3 Description : Artificial Sequence: CEA binding
     r lypupti is
Asy ing Val Cys vid the 1.6 Lys Ash Gin Trp Met Cys Ash Val Let
+111 + +1
+111 + 14
+112 + 147
Full Franchista Native greater
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to its leading the out Artificial Sequences SEA binding
      for a figure of the second
 As in Val. to Ast less the Lys Ash Gin Trp Phe Cys Asp Ala Les
                                     16
 ·2112 16
 47.12. PPT
 <2003> Artificial Sequence
 61.1 22
 <223> Description of Artificial Sequence: CEA binding
      polypeptide
 <400> 82
 Asp Trp Val Cys Gin Tyr Glu Lys Asp Gln Trp Ser Cys Asn Ile Leu
                                     10
 4210> 83
 <211> 16
 <212> PRT
 <213> Artificial Sequence
4.229>
 <223> Description of Artificial Sequence: CEA binding
      polypeptide
r4 U> 83
Asp Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Thr Leu
       - 5
                                    10
<2160 84
<2111 16
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: CEA binding
     polypeptide
Ast Trp Tyr Cys Ash Leu Phe Lys Ash Gln Trp Phe Cys Asp Va. Tyr
           72112 16
<2125 FFT
<917> Artificial Sequence
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ester 1988 mag til bli i flærtidi Stati September (HÆ kindina
    1 2 2 1 2 1 2 2 2 2
 or Tep Tree Cyc. Adm. 1 . The Lyc. According Trp The Cyc. For Fig. 1.4
- lii - ist
- Mi- Artiitilai Sequence
Plans Despriction of Artificial Sequence: CHA binding
     polyses ide
Asy inp Val Cys Oli Pho Pho Lys Lys Gin Trp Phe Cys Ash Leu Leu
                                    1.0
.51% 67
.51% 16
.512% PRT
x213@ Artificial Sequence
ki. 3: Description of Artificial Sequence: CEA binding
     polymentide
< : ... 37
Ash Top Pal Cys Asp Val Leu Lys Trp Gln Trp Pro Cys Asn Ser Tyr
8,115 48
<2112 16
<212> PRT
<2133 Artificial Sequence
Kazaw Description of Artificial Sequence: CEA binding
    relypertible
The Dry Mal Mys Will Dyr wee Lys Diy Win Trp His Dys Ash Lie Let.
Police Artiflical Jeguence
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Part for the training in the Artific transforgular grant CEA binding
       Ang life like the Arm Laudine Lyr Arm Him Try Phot Cys Glin Win Hilk
  1.11 - 16
  · Line FRI
  #// Artificial Sequence
  CE255
  <223> Description of Artificial Sequence: CEA binding
       polypeptide
  <400> 90
  Asp Trp Val Cys Asn Trp Leu Trp Gly Gln Trp Thr Cys Asn Leu Leu
 <210> 91
 <211: 16
 <2121 PRT
 <213> Artificial Sequence
 <220×
 <223> Description of Artificial Sequence: CEA binding
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 <400> 91
 Asp Trp Val Cys Slu Met Phe Lys Lys Gln Trp Val Cys Asn Fro Leu
                        10
 <210> 92
 <211> 16
 <212> PRT
 <213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: CEA binding
      polypeptide
r4302 92
Asp irp lie Cys Ash Teu Phe Lys Ash Gin Trp Phe Cys Gly Pro Leu
· Alz. PRT
*1135 Artificial Sequence
43230 Lescription or Artificial Sequence: CEA binding
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Wig lim Ma. Typ A. Wa. Lie Lyd Arg And Erp Ma. Cyp. Arm Her De.
1.3 / 1:
100 - 14
este Artificial Papir e
+xx > Description of Artificial Sequence: CEA binding
     polypeptide
24005 94
Asr Tip Val Cys Glu Asr. Lys Asr. Phe Lys Trp Phe Cys Asr. Leu Leu
 210 1 90
- 2112 16
<2125 PRT
*113> Artificial Sequence
*2003 - Description of Artificial Sequence: CEA binding
      gblypertide
F : 152 95
Ney Trp Val Cys Glu Tyr Ala Lys Asn Gln Trp Asn Cys Asn Pro 100
                          10
...: 6 . 96
#1.11 / 1k
9..12. HIT
7,139 Artifimial Dequence
<223> Description of Artificial Sequence: CEA binding
      pelypeptide
 Non-lep Tal ly. Ash low involves Ash Gln Trp Phe Cys Glu Trp Ala
                                       1 . .
+..1.+ #7
+..1.+ 1#
+..1.+ ###
+..1++ Artificial #7.px n *2
e Dew Testrightlin in Artificial Gequence: CHA binging
      rc.ypophi m
```

4 1 4

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Non-lig Val. To Alg. Dyn Try Dyn Ern Ein Dry Ene Tys Asin Der Dyn
killsk Artistedal Mequence
<223> Description of Artificial Sequence: CEA binding
      polypeptide
 Asp Irp Tyr Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Leu Val
                      10
2010 - 39
<211 · 16
<212 - PRT
8213 · Artificial Sequence
4220.
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Asn Trp Val Cys Asn Leu Phe Lys Asn Gln Trp Phe Cys Asp Glu Met
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<223> Description of Artificial Sequence: CEA binding
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Asp Trn Val Cys Slu Leu Phe Lys Pro Gln Trp Phe Cys Asn Ile Leu
**11: 16
-212- PRT
KATE: Artificial Sequence
12272
<223> Description of Artificial Sequence: CEA binding
     pulymentide
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Ag in Talliye alife her by Mer all The Ner Tym Act Alaite.
  ester Artists seas segansia
 2223 Description of Artificial Sequence: CEA binding
       polypeptide
 ₹456≥ 102
 Asp Trp Val Cys Asp Tyr Lys Phe Phe Gln Trp Thr Cys Asn Leu Leu
 32105 369
 82112 1F
 <212> PRT
 <213> Artificial Sequence
 <223 Description of Artificial Sequence: CEA binding
       polypeptide
 <400 - 103
 Asr. Orp Val Cys Glu Trp Leu Lys Pro Gln Trp Trp Cys Asn Ser Tyr
                                     10
<210: 104
<2110 16
<2120 PRT
<213> Artificial Sequence
₹2295
<223> Description of Artificial Sequence: CEA binding
      polypeptide
<400> 104
Aso Trp Val Cys Glu Phe Phe Lys Pro Gln Trp Met Cys Asn Ile Leu
· .11. · 14
**12 FRI
Philip Artificial Sequence
. 42.00
<223> Description of Artificial Sequence: CEA binding
     polypeptide
44112 115
```

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Activity

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**Comparison of Artificial Sequence: CEA binding pol
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